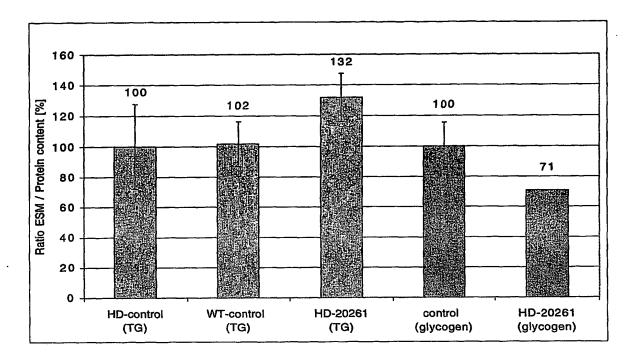
Figure 1. Energy storage metabolite content of a Drosophila PRL-1 mutant



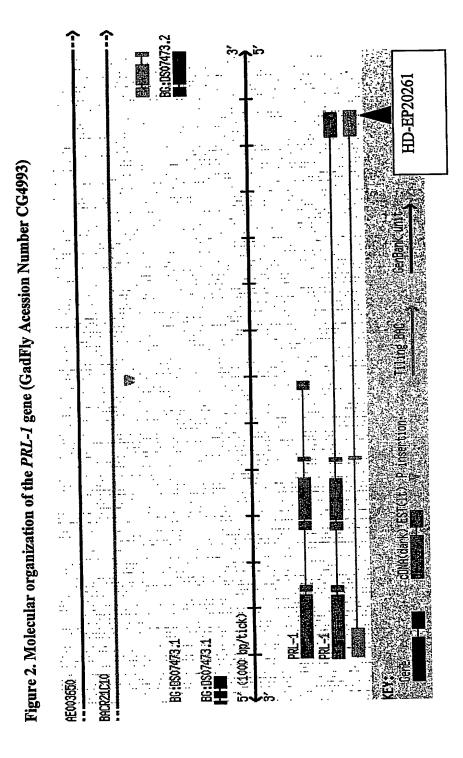


Figure 3. Nucleic acid sequences encoding the human proteins of the invention and amino acid sequences of the proteins of the invention

Figure 3A. Homo sapiens protein tyrosine phosphatase type IVA, member 1 (Prl-1), Nucleic acid sequence (SEQ ID NO: 1)

	_					
1	ccggctcggt	acgcgctctg	ctccgagccg	ctcactgcgt	ggtagagtct	ggtgccccg
61	contraceta	categeegee	accoccoctc	cgccacgacc	accgccgccc	eetgeeetge
121	SACCSCCACC	accacctata	tcaccaccac	ctcgggaccg	gctgtatgat	caggecacaa
1 2 1	tetteaatga	gtaaacatat	tcctcaattc	tgtggtgttc	ttggtcacac	attlatygag
2/1	tttctgaagg	gcagtggaga	ttactgccag	gcacagcacg	acctctatgc	agacaagtga
201	actotagaaa	ctgattactg	ctccaccaag	aagcccccat	aagagtggtt	atcctggaca
261	cacaactett	gaattgaaat	ccacagagca	ttttacaaga	gttctgacct	ggatggggta
421	aacctcagtg	cacttetttt	ctattaacct	cagtattact	ggattgaaga	attgetgett
101	cttattagga	ggttcatttc	acttatcatt	acttacaact	tcatactcaa	agcactgaga
5/1	atttcaagtg	gagtatattg	aagtagactt	cagtttcttt	gcatcatttc	tgtattcaat
601	++++++aatt	atttcataac	cctattgagt	gttttttaac	taaattaaca	tggcicgaal
661	assccacces.	actectataa	aagtcacata	caagaacatg	agatttctta	ttacacacaa
721	tccaaccaat	gcgaccttaa	acaaatttat	agaggaactt	aagaagtatg	gagttaccac
791	aataqtaaqa	gtatgtgaag	caacttatga	cactactctt	gtggagaaag	aaggtaccca
241	tattattaat	taacctttta	atgatggtgc	accaccatcc	aaccagattg	ttgatgattg
9.01	gttaagtett	gtgaaaatta	agtttcgtga	agaacctggt	tgttgtattg	etgeteatty
961	cattacadac	cttgggagag	ctccagtact	tgttgcccta	gcattaattg	aaggtggaat
1021	gaaatacgaa	gatgcagtac	aattcataag	acaaaagcgg	cgtggagctt	ttaacagcaa
1081	gcaacttctg	tatttggaga	agtatcgtcc	taaaatgcgg	ctgcgtttca	aagatteeaa
1141	cootcataga	aacaactott	gcattcaata	aaattggggt	gcctaatgct	actggaagtg
1201	gaacttgaga	tagggcctaa	tttgttatac	atattagcca	acatgttggc	ttagtaagte
1261	taatgaagct	tccataggag	tattgaaagg	cagttttacc	aggcctcaag	ctagacagat
1321	ttggcaacct	ctgtatttgg	gttacagtca	acctatttgg	atacttggca	aaagattett
1381	gctgtcagca	tataaaatgt	gcttgtcatt	tgtatcaatt	gacettteee	caaaicatge
1441	agtattgagt	tatgacttgt	taaatctatt	cccatgccag	aatcttatca	atacataaya
1501	aatttaggaa	gattaggtgc	caaaataccc	agcacaatac	ttgtatattt	tagtaccac
1561	acagaagtaa	aatcccagga	actatgaaca	ctagacetta	tgtggtttat	statttagst
1621	atttcaaaca	ttgaaagtag	ggcctacatg	gttatttgcc	tgeteaettt	atytttacat +++++++
1681	ctcccacatt	cataccaata	tacgtcaggt	ttgcttaacc	actgattttt	~~*
1741	accaagtctt	acagtgatta	ttttacgtgt	ttccatgtat	ctcactttgt	tetattasas
1801	aaaaacctcc	attttgaaaa	tctacgttgt	acagaagcac	atgiciliaa	caccaga
1861	caaaaaagcc	ttacattaat	ttaatgtttg	cactctgagg	tgcaacttaa	cagggagggc
1921	ctgagaaaag	aatgggaggg	ggctattaat	tatttttag	caaaatgttg	nggtagagat
1981	gtgcaaacat	gtagaatatg	ctctttaatt	tagtaaaata	-ttttttaaa	aggragagar
2041	gctttgttat	tgtaatcata	aacttcctga	aattettgta	. accentices	catacctacc
2101	. agaagtgtgt	ttaccaactt	atttttgttt	gaaagtgtga		astatttat
2161	. ctctcttgca	aaaaaagaaa	tgggtttctg	ctaatgaatt	gageaeatet	tattatataa
2221	atgccttttg	gagctgtgta	agttaatatt	: tgatacttga	caatttgttt	tatatagaaa
2281	ttgataaaat	ggtgatgtgt	attaatgtta	gtteaaceat	atattiatac	ttataaaaac
2341	tgtgtggtta	. tagttctgtg	ggagaaataa	ttttgtcagt	. gillaccago	cattracatt
2401	ttagtgcgag	agctgaaaca	tctaaataa	taatgacatg	catteaceat	cactgagact
2461	ggtttgcttt	aaattaactt	attttgtaga	agacaaaatg	aattycact	ttttagtaaa
2521	gtgtcctcat	ctttttacaa	ataaatgaag	gattataaat	. gatyttagta	ttttagtaaa
2583	L cttttagaca	aaatttgtta	gggtcattca	. tgaaaacctt	. dalautaada	gcactttcca
2641	L ttatatactt	tttaaaggto	tagataatt	. cgaaccaatt	. cartattyty	tactgaggag
270	L aaataatgta	ı tagtagagga	cagccttggt	tegtaaaget	. cagilledaci	agttcatggt
2761	L tttgtgcaac	: ttctgagcct	cagttttctc	ctttgcaaat	aa.aa.laC	atacctttat
282:	ı agattttgaa	attaatttaa	atattagta	ciggtacate	, aayyuuaat . aastotaost	gttaagtttc
2883	L ctttaatgat	ccacaataat	ccctttgate	acgutaaccu	. aaatttaagat	gtctttgtct
294:	. aattttttt	gaatagcagt	cataaatgta	a aayyactcaa	agullaagu	aaaagtgata
300:	l ctccaccttg	, tgtttcaaag	g aatttagtto	cacetette	Lactagette	acacttaata

```
3061 tatttcattg gattttagac agggcaaaag gaagaacagg ggcctctgga ggcccttggt
3121 tatttaaatc ttggattatt tgtgatagta atcacaaatt tttggctaat ttttaacctg
3181 aggttttgtt tttttttaa aggaaatgca gcctagtctt gagaacataa ttttatataa
3241 tcaattacta aatgttaaac tattaccaca cagcccataa aacagcattt gcgtttattg
3301 agagagaga tgtgccatca tgattaatga aaactatctt ttgagtttga aaagaaatta
3361 atttgcagtg tttggattgt atatatggtg ctaaaaataa attaatttac tttataaacc
3421 ttatctgtac attatacgat gtgatgaaat ttgcttttta tccaaatatt ttgtatcttg
3481 taaatatggc taattatagg aatgcctata atacatctta gattccttat atctaataag
3541 agttcaaaga gttatgagtt gaagtcttga atgcaggaaa ctatctgata gtgttctaaa
3601 atttggttac ttgggtttgg ataccettag tgggatgatg taaatagagg ctagetacet
3661 aggettgtet atageaacca taatgttgat gtaagtaatg eggttaetga ateataagaa
3721 aatgccatct ctttttagtt gaaggaaaac tctggaagta ggtgccattg gtcattctgc
3781 agtgcactgc aaccattgtt tcccctagtg ccctcttttc cctagggcat tgctctccta
3841 ttcccacgcc ttaacacagc tctataccta gaagcagcca gcccaggcat gcagtcacat
3901 ttaatcacat ccccttcta gagtgcttca aaatgatgta gtccctcaac ttggctaaag
3961 aatctcaatc tcttgaaatt tatttttta atgtcatatt catctggtaa atatctactg
4021 tttgccaggc atttaagaat atggcaaaga acataaaaga tggtgtcacc agattttggt
4081 caccaatgag tacccgaccc gttgccatga ttaagagaga atgctttcta ttggagtttc
4141 aggaaatata atttgagaat actttaaagg gaagtggaag tataagtgaa tgatattttt
4261 agtgtctgcc atacatgtta atattctaca ttcttgcttc cttaaattaa tatgtttgtg
4321 tgtatatatg tgcctcacac ctgaattgaa aattaaagac tggtttaaaa gtggttaaaa
4381 aaaaaaaaaa aaaa
```

Figure 3B. Homo sapiens protein tyrosine phosphatase type IVA, member 1 (Prl-1), Amino acid sequence (SEQ ID NO: 2)

```
1 marmnrpapv evtyknmrfl ithnptnatl nkfieelkky gvttivrvce atydttlvek
61 egihvldwpf ddgappsnqi vddwlslvki kfreepgcci avhcvaglgr apvlvalali
121 eggmkyedav qfirqkrrga fnskqllyle kyrpkmrlrf kdsnghrnnc ciq
```

Figure 3C. Homo sapiens protein tyrosine phosphatase type IVA, member 2 (Prl-2), Nucleic acid sequence, transcript variant 1 (SEQ ID NO: 3)

```
1 ageggggetg egegaagtea tegetgttee agacagegat gaetegagag eggtgggggt
 61 ggcggcgcga tcggccgggc tgtaaccgtc gtctgtccgg gagcggctgg agcggcagcg
121 geggeeggge aeggegegag gtgaegeeae agggeagegg eggeagegga ggeageggeg
181 gcagcaggag acgcagcggc ggccgcagca gcagcagcaa gacggactcg tggagacgcg
241 ccgccgccgc cgccgccggg ccgggccggg tgtcgcgcgc cgaggctggg ggggagtcgt
301 cgccgccgcc gccaccgcta ccgccgccgc cgccgccgcc gaggtgactg aggagagagg
361 cgcctcctcg ctcccgccac cgccggactt caatgcccag tccccagctc gccagcgttt
421 ttcgttggaa tatacgttgc acatttatgg cgattctgag tgtgagggca gacttctgcc
481 aggeteagea cageatttte getgacaagt gagettggag gttetatgtg ceataattaa
541 cattgccttg aagactcctg gacaccgaga ctggcctcag aaatagttgg ctttttttt
601 tttttaattg caagcatatt tcttttaatg actccagtaa aattaagcat caagtaaaca
661 agtggaaagt gacctacact tttaacttgt ctcactagtg cctaaatgta gtaaaggctg
721 cttaagtttt gtatgtagtt ggattttttg gagtccgaat atttccatct gcagaaattg
781 aggcccaaat tgaatttgga ttcaagtgga ttctaaatac tttgcttatc ttgaagagag
841 aagcttcata aggaataaac aagttgaata gagaaaacac tgattgataa taggcatttt
901 agtggtcttt ttaatgtttt ctgctgtgaa acatttcaag atttattgat tttttttt
961 cactttcccc atcacactca cacgcacgct cacacttttt atttgccata atgaaccgtc
1021 cagcccctgt ggagatctcc tatgagaaca tgcgttttct gataactcac aaccctacca
1081 atgctactct caacaagttc acagaggaac ttaagaagta tggagtgacg actttggttc
1141 gagtttgtga tgctacatat gataaagctc cagttgaaaa agaaggaatc cacgttctag
1201 attggccatt tgatgatgga gctccacccc ctaatcagat agtagatgat tggttaaacc
```

1261	tgttaaaaac	caaatttcgt	gaagagccag	gttgctgtgt	tgcagtgcat	tgtgttgcag
4 2 2 4		~~~~~~tata	CTCCTCCCC	ttactitaat	Lyaalylgga	acgaag
4004		+~~~+++>+>	araraaaaaa	daaddddadc	u L L Caa L L C C	auacagoogo
4 4 4 4		assatarcas.	cctaagatgc	gattacgett	cagagatace	aacgggcacc
4 - 01		at agaaggaa	arotaaacoa	aggetgaett	gacegeea	40005055
		aataassta	traatctroa	atattacctu	Lyteattaaa	gcagcgacgg
4		aatcaaccac	tctcctaato	attogaacaa	aaycaaacaa	aaaagaaa
		~+~~ <u>~</u> +~~~	tatttaaaaa	aacacaaaca	yaaaayyaac	caacccag
		++~~+~~+	ttttaaaatt.	roaatttctu	Ccayyaccya	accaccega
		a+++++>>>	tttttcaaaa	Eaggicula	ayyaaaacca	gcagaacar
	· · · · · · · · · · · · · · · · · · ·		attrananaa	cacactcttc	Cattatyctc	ggowows
		~~~~~~++++	TETETECCCC.	EEEEEEGLAA	gggagggccg	gcggcacac
4001	aatat	+++++~~++~	ctctcctaca	TCTCCCTLLL	CCCCGaccc	aageegeaga
0044	+t-a	accettatta	ctgtagatgt	gcgtgcagtc	Lygcaycece	aagcccacc
-4-04		~~~+~~~~~	222222222C	aaaaaacaac	accaaaaaaa	cagcagogae
2101	atatatattc	agacadadada	ttagtcttta	ctgatgaaag	ggtgttcatg	ttagtttctt
		+~+>>+>	aacaaaataa	ccaadadcci	LLLyLLLLy	CCCCCCCCC
			cattttaaca	agagecee	Cicaacicac	ccgagagaag
2281	ataaattagt aattcttctc	ttagataacgg	aatgaaggta	agtggttatc	ccagaaactt	gtcttctaaa
			2242227277	TCCACCCACL	daucutacce	CCCGGGGGGG
2401	gtagaattgt	ceaggecate	attactcta	ttttctatct	agtatcagag	aatgctggta
2461	gcagaactgc gcttaaaatt	gggctgtagc	gettetact	ctgaattttc	aggaaccgtc	aaaggagcag
2521	cagcaaattc		gacttgaga	atacttataa	tatgtgtttt	ccaaactgcc
2581	cagcadatic	acacacccc	ttaaccacto	attoccttot	tattactagg	ttttttgaga
2641	. ccctatatgt . ttaaaaaaaa	aaagttcagt	otttaaaacc	aacaatgatg	cctagtgagt	atgtgtccac
		~~~~+ ~~~~	adadacarco	Tocaacccaa	Luagtagtgu	uggguvugug
2761	aggecataac	agggcagaag	tagagacatttt	gegattett	aactaaattt	agtgtactga
2821	. ttgcttgtga	ageggigtag	caycatttat	goagaceart	tatgatcagg	ctgcatggac
2881	. tctagaaaag	ctgtttttt	atcacccccqc	cttocactat	tttcacctct	aaatattacg
2941	aaagcaggta	gaggggcacc	attaggggtt	. daatacqqq	ttaaagtcat	cttgtcctgc
3001	tactcagtag	tgeeetgett	. ccagggeeee	. gaatatgggt	agcgtcagct	aggccaatag
3063	tggaatttgc	: tgtgcagagc	talaaguut	. codecetgee	atottoacco	gctatgtgaa
312:	L gaacagaccg	ggaccttgtc	: ccacaccyat	. tttaactaaa	cocaaatcto	tagattetet
3183	L ctgcctattt	cctatgctgg	agittigati	. cctaactaaa	gaaattottt	ctaggacttt
324:	L ceteteceat	cccagaaaac	addacadaa	. tttgagaato	caacaataga	ttccattaat
330:	l aaaacataat	ggtatatcca	aaattetti	taagaacy	tagatggtgt	ttccattaat
336	l atagactcaa	a gatcaaaaca	geatacety	: taayctaage	: cagacggogo	tgattccact
342	l gggttttgat	caatacaata	acaaacctt	t-tatatatat	. atactotgus	ttttgttgtt tatatatata
348	l tggggggagg	g gggtgtgtgt	grararata	gigigigigigi	taggaaaatt	tgtgtgtgtg
354	l tgcacgcgca	a gtgtccatca	a gratcagrad	- pagatassa	: attacccto	acattcctgg
360	1 ttctgtattq	g aggagaagga	tgtataaag	- aacacyadac	· ccttaaagca	cttttattt
366	1 aaagactaa	t gttaattgt!	cttaaaact	g yarrritt	. catamamam	a atttttttct
372	1 tttcgattta	a atgaagtati	geragerga	a godayttige	a accettta	tgtcagattg
378	1 atttgaaag	g tgtgcagcci	c gatttaaaa	c caaaccctg	2 22222222	a agaacaataa
384	1 aacatattt	t acacgctca	a aaaaaaaaa	a aaaaaadaa	a aaaaaaaaaaa	a aaaaaaaaaa
390	1 aaaaaaaaa	a aaaaaaaaa	a aaaaa			

Figure 3D. Homo sapiens protein tyrosine phosphatase type IVA, member 2 (Prl-2), Amino acid sequence transcript variant 1 (SEQ ID NO: 4)

```
1 mnrpapveis yenmrflith nptnatlnkf teelkkygvt tlvrvcdaty dkapvekegi
61 hvldwpfddg apppnqivdd wlnllktkfr eepgccvavh cvaglgrapv lvalaliecg
121 mkyedavqfi rqkrrgafns kqllylekyr pkmrlrfrdt nghccvq
```

Figure 3E. Homo sapiens protein tyrosine phosphatase type IVA, member 2 (Prl-2), Nucleic acid sequence, transcript variant 2 (SEQ ID NO: 5)

```
1 agcggggctg cgcgaagtca tcgctgttcc agacagcgat gactcgagag cggtgggggt
 61 ggcggcgcga tcggccgggc tgtaaccgtc gtctgtccgg gagcggctgg agcggcagcg
121 geggeegge aeggegegag gtgaegeeae agggeagegg eggeagegga ggeageggeg
181 gcagcaggag acgcagcggc ggccgcagca gcagcagcaa gacggactcg tggagacgcg
241 ccgccgccgc cgccgcggg ccgggccggg tgtcgcgcgc cgaggctggg ggggagtcgt
301 egeogeegee gecacegeta eegeogeege egeogeege gaggtgaetg aggagagagg
361 egecteeteg etceegecac egeeggactt caatgeceag teceeagete geeagegttt
421 tregttggaa tatacgttgc acatttatgg cgattctgag tgtgagggca gacttctgcc
481 aggctcagca cagcattttc gctgacaagt gagcttggag gttctatgtg ccataattaa
541 cattgccttg aagactcctg gacaccgaga ctggcctcag aaatagttgg ctttttttt
601 tttttaattg caagcatatt tcttttaatg actccagtaa aattaagcat caagtaaaca
661 agtggaaagt gacctacact tttaacttgt ctcactagtg cctaaatgta gtaaaggctg
721 cttaagtttt gtatgtagtt ggattttttg gagtccgaat atttccatct gcagaaattg
781 aggcccaaat tgaatttgga ttcaagtgga ttctaaatac tttgcttatc ttgaagagag
841 aagcttcata aggaataaac aagttgaata gagaaaacac tgattgataa taggcatttt
901 agtggtcttt ttaatgtttt ctgctgtgaa acatttcaag atttattgat tttttttt
961 cactttcccc atcacactca cacgcacgct cacacttttt atttgccata atgaaccgtc
1021 cageceetgt ggagatetee tatgagaaca tgegttttet gataaeteae aaceetaeea
1081 atgctactct caacaagttc acagaggaac ttaagaagta tggagtgacg actttggttc
1141 gagtttgtga tgctacatat gataaagctc cagttgaaaa agaaggaatc cacgttctag
1201 attggccatt tgatgatgga gctccaccc ctaatcagat agtagatgat tggttaaacc
1261 tgttaaaaac caaatttcgt gaagagccag gttgctgtgt tgcagtgcat tgtgttgcag
1321 gattgggaag ggcacctgtg ctggttgcac ttgctttgat tgaatgtgga atgaagtacg
1381 aagatgcagt tcagtttata agacaaaaaa gaaggggagc gttcaattcc aaacagctgc
1441 tttatttgga gaaataccga cctaagatgc gattacgctt cagagatacc aatgggcatt
1501 gctgtgttca gtagaaggaa atgtaaacga aggctgactt gattgtgcca tttagaggga
1561 actottggta cotggaaatg tgaatotgga atattacotg tgtcatcaaa gtagtgatgg
1621 attcagtact cctcaaccac tctcctaatg attggaacaa aagcaaacaa aaaagaaatc
1681 tetetataaa atgaataaaa tgtttaagaa aagagaaaga gaaaaggaat taatteagtg
1741 aaggatgatt ttgctcctag ttttggagtt tgaatttctg ccaggattga attattttga
1801 aatctcctgt ctttttaaac tttttcaaaa taggtctcta aggaaaacca gcagaacatt
1861 aggcctgtgc aaaaccatct gtttggggag cacactcttg gcacatagat ctccctgtgg
1921 tgggattttt ttcccttttt ttgtggggga gggttggtgg tatatttttc ccctctttt
1981 tectteetet ectacatete cetttteece egatecaagt tgtagatgga atagaageee
2041 ttgttgctgt agatgtgcgt gcagtctggc agccttaagc ccac
```

Figure 3F. Homo sapiens protein tyrosine phosphatase type IVA, member 2 (Prl-2), Amino acid sequence transcript variant 2 (SEQ ID NO: 6)

```
1 mnrpapveis yenmrflith nptnatlnkf teelkkygvt tlvrvcdaty dkapvekegi
61 hvldwpfddg apppnqivdd wlnllktkfr eepgccvavh cvaglgrapv lvalaliecg
121 mkyedavqfi rqkrrgafns kqllylekyr pkmrlrfrdt nghccvq
```

Figure 3G. Homo sapiens protein tyrosine phosphatase type IVA, member 2 (Prl-2), Nucleic acid sequence, transcript variant 3 (SEQ ID NO: 7)

```
1 ageggggetg egegaagtea tegetgttee agacagegat gaetegagag eggtgggggt
61 ggeggegega teggeeggge tgtaacegte gtetgteegg gageggetgg ageggeageg
121 geggeeggge aeggegegag gtgaegeeae agggeagegg eggeagegga ggeageggeg
181 geageaggag aegeagegge ggeegeagea geageageaa gaeggaeteg tggagaegeg
```

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241 ccgccgccgc cgccgccggg ccgggccggg tgtcgcgcgc cgaggctggg ggggagtcgt
301 cgccgccgcc gccaccgcta ccgccgccgc cgccgccgcc gaggtgactg aggagagag
361 cgcctcctcg ctcccgccac cgccggactt caatgcccag tccccagctc gccagcgttt
421 tregttggaa tatacgttgc acatttatgg cgattctgag tgtgagggca gacttctgcc
481 aggctcagca cagcattttc gctgacaagt gagcttggag gttctatgtg ccataattaa
541 cattgccttg aagactcctg gacaccgaga ctggcctcag aaatagttgg ctttttttt
601 tttttaattg caagcatatt tcttttaatg actccagtaa aattaagcat caagtaaaca
661 agtggaaagt gacctacact tttaacttgt ctcactagtg cctaaatgta gtaaaggctg
721 cttaagtttt gtatgtagtt ggattttttg gagtccgaat atttccatct gcagaaattg
781 aggcccaaat tgaatttgga ttcaagtgga ttctaaatac tttgcttatc ttgaagagag
841 aagcttcata aggaataaac aagttgaata gagaaaacac tgattgataa taggcatttt
901 agtggtcttt ttaatgtttt ctgctgtgaa acatttcaag atttattgat tttttttt
961 cactttcccc atcacactca cacgcacgct cacacttttt atttgccata atgaaccgtc
1021 cagcccctgt ggagatetee tatgagaaca tgcgttttet gataacteac aaccctacca
1081 atgctactct caacaagttc acagaggaac ttaagaagta tggagtgacg actttggttc
1141 gagtttgtga tgctacatat gataaagctc cagttgaaaa agaaggaatc cacgttctaa
1201 aaaagaaggg gagcgttcaa ttccaaacag ctgctttatt tggagaaata ccgacctaag
1261 atgcgattac gcttcagaga taccaatggg cattgctgtg ttcagtagaa ggaaatgtaa
1321 acgaaggetg acttgattgt gccatttaga gggaactett ggtacetgga aatgtgaate
1381 tggaatatta cctgtgtcat caaagtagtg atggattcag tactcctcaa ccactctcct
1441 aatgattgga acaaaagcaa acaaaaaaga aatctctcta taaaatgaat aaaatgttta
1501 agaaaagaga aagagaaaag gaattaattc agtgaaggat gattttgctc ctagttttgg
1561 agtttgaatt tctgccagga ttgaattatt ttgaaatctc ctgtcttttt aaactttttc
1621 aaaataggtc tctaaggaaa accagcagaa cattaggcct gtgcaaaacc atctgtttgg
1681 ggagcacact ctt
```

Figure 3H. Homo sapiens protein tyrosine phosphatase type IVA, member 2 (Prl-2), Amino acid sequence, transcript variant 3 (SEQ ID NO: 8)

1 mnrpapveis yenmrflith nptnatlnkf teelkkygvt tlvrvcdaty dkapvekegi 61 hvlkkkgsvq fqtaalfgei pt

Figure 3I. Homo sapiens protein tyrosine phosphatase type IVA, member 3 (Prl-3), Nucleic acid sequence (SEQ ID NO: 9)

```
61 gggggggggg cgggctgttt tgttcctttt cttttttaag agttgggttt tctttttaa
121 ttatccaaac agtgggcagc ttcctcccc acacccaagt atttgcacaa tatttgtgcg
181 gggtatgggg gtgggttttt aaatctcgtt tctcttggac aagcacaggg atctcgttct
241 ceteattttt tgggggtgtg tggggactte teaggtegtg tececageet tetetgeagt
301 ceettetgee etgeegggee egtegggagg egceatgget eggatgaace geeeggeece
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421 gctcagcacc ttcattgagg acctgaagaa gtacggggct accactgtgg tgcgtgtgtg
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541 gtttgacgat ggggcgcccc cgcccggcaa ggtagtggaa gactggctga gcctggtgaa
601 ggccaagttc tgtgaggccc ccggcagctg cgtggctgtg cactgcgtgg cgggcctggg
661 ccgggctcca gtccttgtgg cgctggcgct tattgagagc gggatgaagt acgaggacgc
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841 gtgctgcgtt atgtagctca ggaccttggc tgggcctggt cgtcatgtag gtcaggacct
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1021 gcgaggagcc cctcgggccc tgggtggcct ctgggccctt tctcctgtct ccgccactcc
1081 ctctggcggc gctggccgtg gctctgtctc tctgaggtgg gtcgggcgcc ctctgcccgc
1141 cccctcccac accagccagg ctggtctcct ctagcctgtt tgttgtgggg tgggggtata
```

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Figure 3J. Homo sapiens protein tyrosine phosphatase type IVA, member 3 (Prl-3), Amino acid sequence (SEQ ID NO: 10)

1 marmnrpapv evsykhmrfl ithnptnatl stfiedlkky gattvvrvce vtydktplek 61 dgitvvdwpf ddgapppgkv vedwlslvka kfceapgscv avhcvaglgr apvlvalali 121 esgmkyedai qfirqkrrga inskqltyle kyrpkqrlrf kdphthktrc cvm

Figure 4. Expression of Prl-1 in different mammalian models

Figure 4A. Real-time PCR analysis of Prl-1 expression in wild type mouse tissues

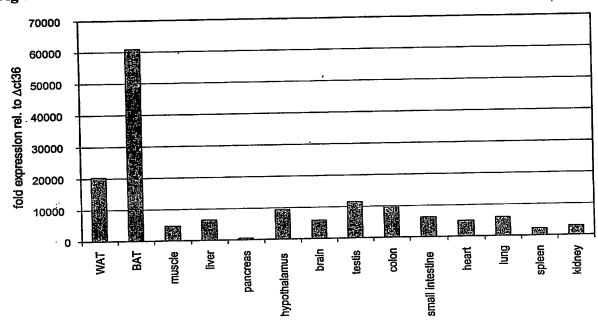


Figure 4B. Real-time PCR analysis of Prl-1 expression in different mouse models

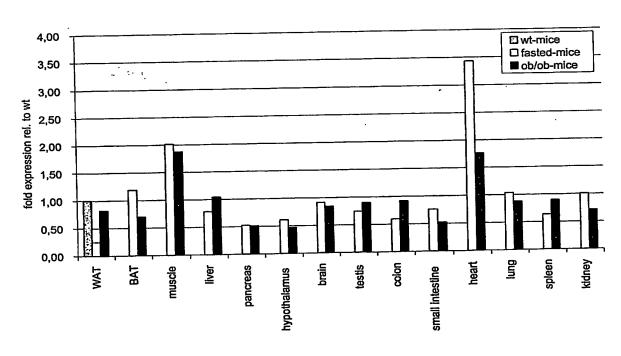


Figure 4C. Real-time PCR analysis of Prl-1 expression in adipocytes during differentiation of 3T3-L1 cells from preadipocytes to mature adipocytes

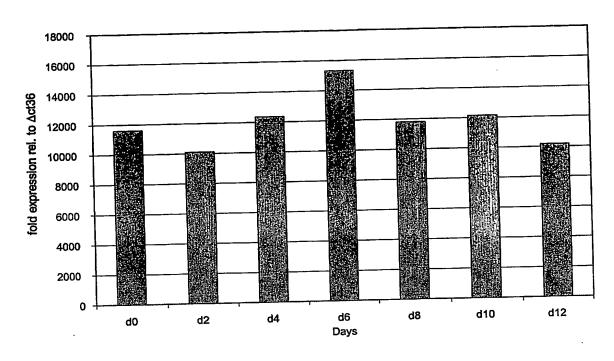


Figure 4D. Real-time PCR analysis of Prl-3 expression in wild type mouse tissues

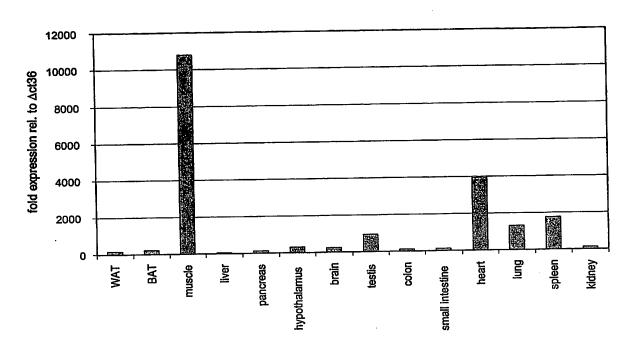
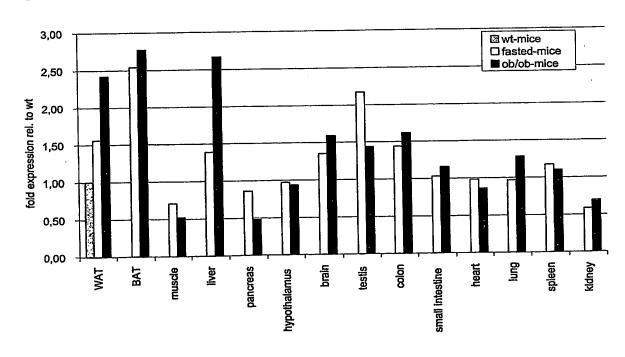


Figure 4E. Real-time PCR analysis of Prl-3 expression in different mouse models



Fgure 4F. Real-time PCR analysis of Prl-3 expression in wild type mice fed a high fat diet compared to mice fed a control diet

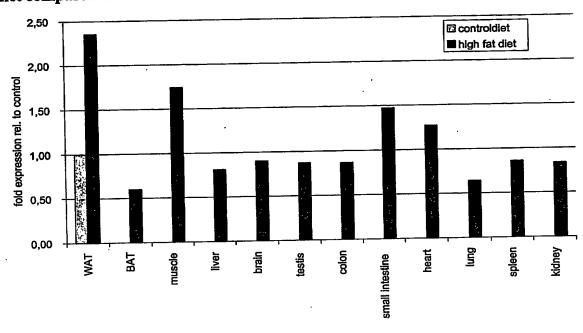


Figure 4G. Real-time PCR analysis of Prl-3 expression in adipocytes during differentiation of 3T3-L1 cells from preadipocytes to mature adipocytes

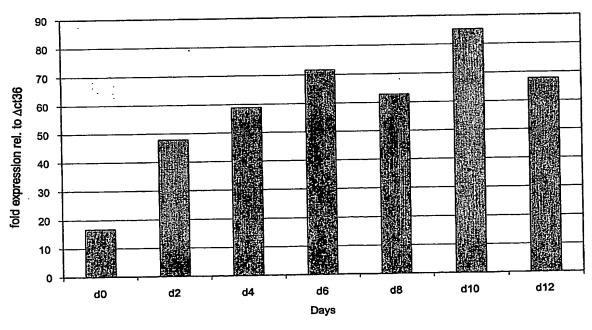


Figure 5. In vitro assays for determination of triglyceride storage and glycogen levels in adipocytes overexpressing Prl-1

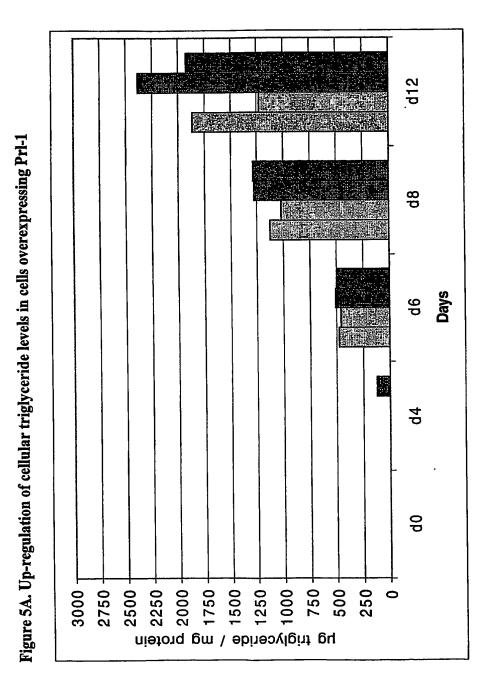


Figure 5B. Up-regulation of cellular glycogen levels in cells overexpressing Prl-1

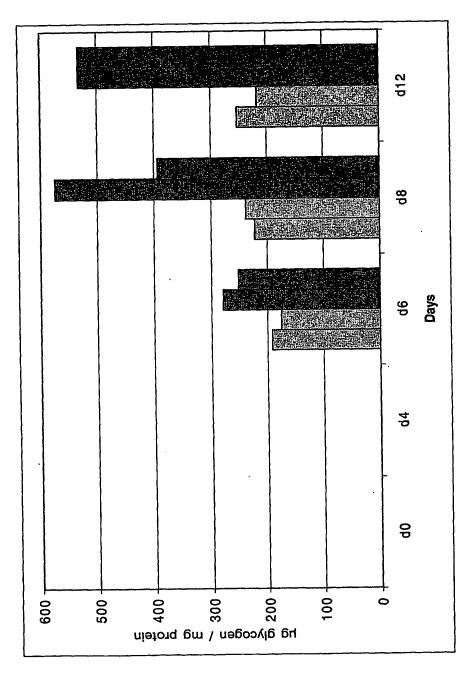


Figure 6. In vitro assays for determination of lipid synthesis and esterification of free fatty acids in Prl-1 loss of function (LOF) adipocytes

Figure 6A. Lipid synthesis levels on day 6 of differentiation in Prl-1 LOF 3T3-L1 cells

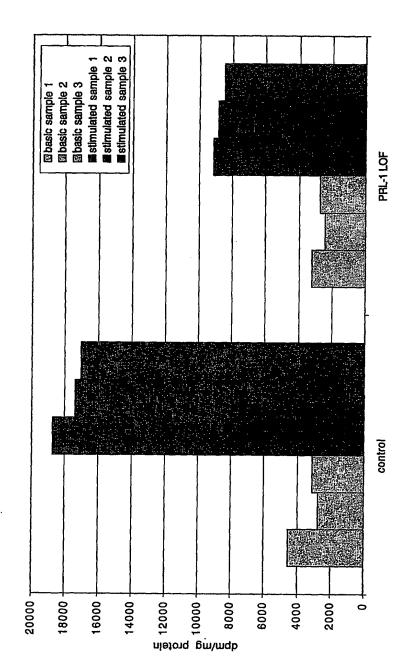


Figure 6B. Fatty acid esterification levels after free fatty acid uptake on day 12 of differentiation in PrI-1 LOF 3T3-L1 cells

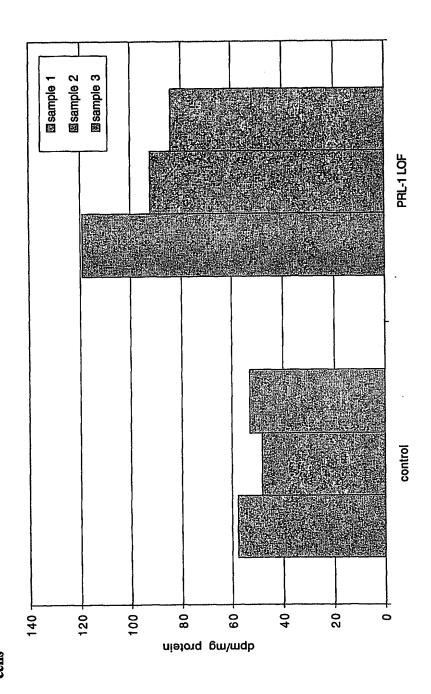


Figure 7. Expression of human PRL-1 homologs in mammalian (human) tissue

Figure 7A. Microarray analysis of PRL-1 expression in abdominal derived primary adipocyte cells during the differentiation from preadipocytes to mature adipocytes

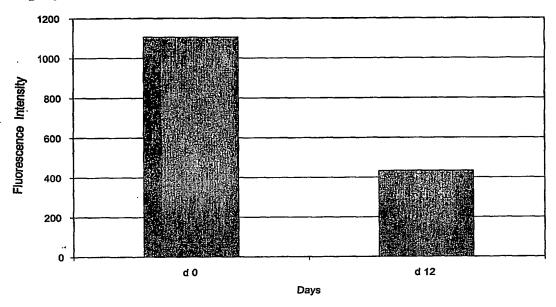


Figure 7B. Microarray analysis of PRL-1 expression in a human adipocyte cell line during the differentiation from preadipocytes to mature adipocytes

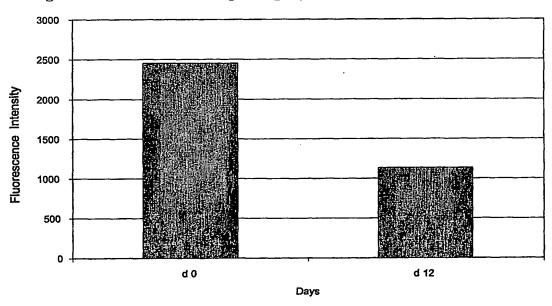


Figure 8. Real-time PCR analysis of the expression of *PRL-1* homologs in different human tissues

Figure 8A. Real-time PCR analysis of PRL-1 expression in different human tissues

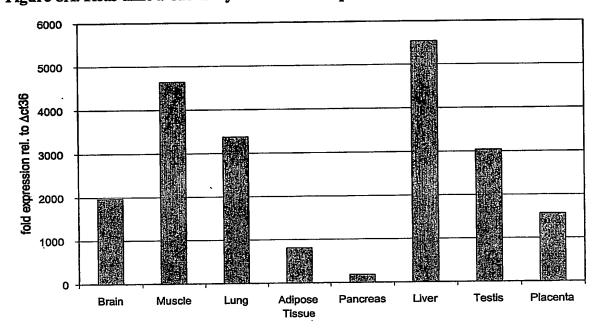


Figure 8B. Real-time PCR analysis of PRL-2 expression in different human tissues

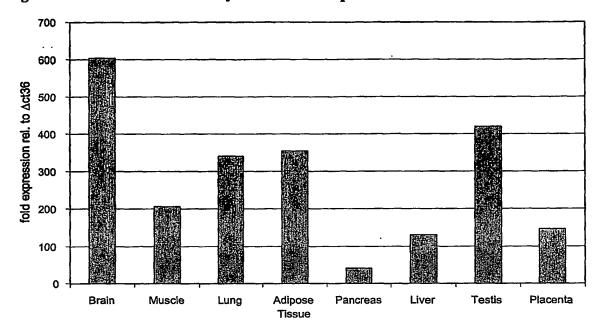
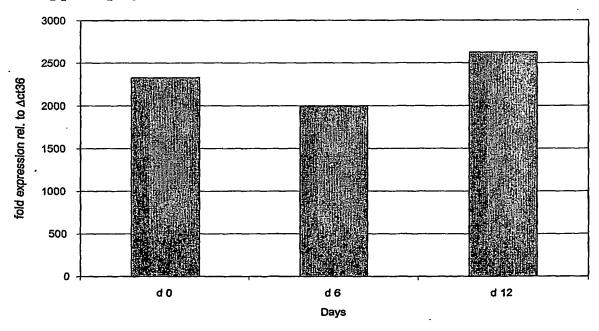


Figure 8C. Real-time PCR analysis of PRL-2 expression in human primary adipocytes during preadipocyte differentiation



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Figure 8D. Real-time PCR analysis of PRL-3 expression in different human tissues

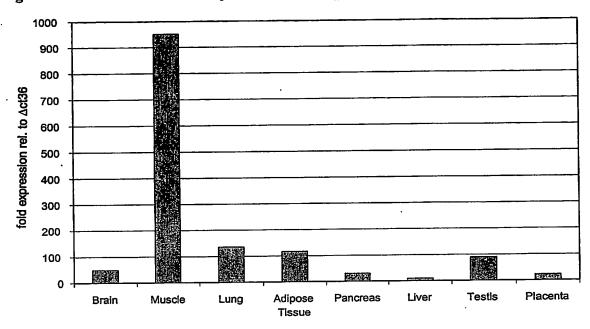


Figure 8E. Real-time PCR analysis of PRL-3 expression in human primary adipocytes during preadipocyte differentiation

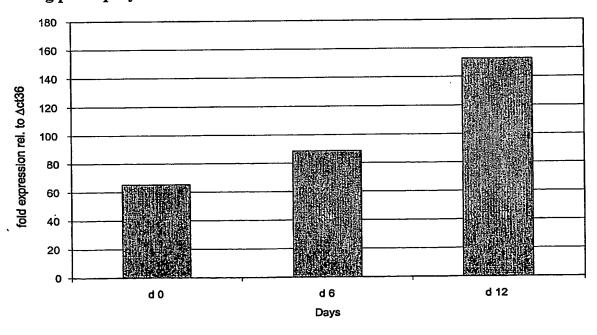


Figure 9. In vitro assays for determination of free fatty acid and glucose uptake by adipocytes overexpressing Prl-1

Figure 9A. Up-regulation of free fatty acid uptake in SGBS cells overexpressing Prl-1

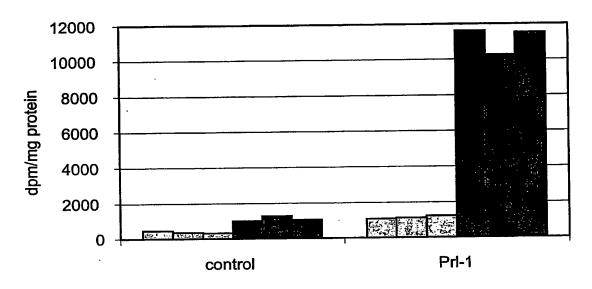


Figure 9B. Up-regulation of glucose uptake in SGBS cells overexpressing Prl-1

